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(d)

CLAIMS

What is claimed is:

- An isolated nucleic acid fragment encoding all or a substantial portion of an Arabidopsis thaliana-like sugar transport protein comprising a member selected from the group consisting of:
 - an isolated nucleic acid fragment encoding all or a substantial portion of the amino acid sequence set forth in a member selected from the group consisting of SEQ ID NO:2, 4, 6, 8, 10, 12, 14 and 16;
 - (b) an isolated nucleic acid fragment that is substantially similar to an isolated nucleic acid fragment encoding all or a substantial portion of the amino acid sequence set forth in a member selected from the group consisting of SEQ ID NO:2, 4, 6, 8, 10, 12, 14 and 16; and
 - an isolated nucleic acid fragment that is complementary to (a) or (b).
- The isolated nucleic acid fragment of Claim 1 wherein the nucleotide sequence of the fragment comprises all or a portion of the sequence set forth in a member selected from the group consisting of SEQ ID NO:1, 3, 5, 7, 9, 11, 13 and 15.
 - 3. A chimeric gene comprising the nucleic acid fragment of Claim 1 operably linked to suitable regulatory sequences.
 - 4. A transformed host cell comprising the chimeric gene of Claim 3.
- An Arabidopsis thaliana-like sugar transport protein polypeptide comprising all or a substantial portion of the amino acid sequence set forth in a member selected from the group consisting of SEQ ID NO:2, 4, 6, 8, 10, 12, 14 and 16.
 - 6. An isolated nucleic acid fragment encoding all or a substantial portion of an Beta vulgaris-like sugar transport protein comprising a member selected from the group consisting of:
 - an isolated nucleic acid fragment encoding all or a substantial portion of (a) the amino acid sequence set forth in a member selected from the group consisting of SEQ ID NO:18, 20, 22, 24, 26 and 28;
 - an isolated nucleic acid fragment that is substantially similar to an (b) isolated nucleic acid fragment encoding all or a substantial portion of the amino acid sequence set forth in a member selected from the group consisting of SEQ ID NO:18, 20, 22, 24, 26 and 28; and
 - an isolated nucleic acid fragment that is complementary to (a) or (b).
- The isolated nucleic acid fragment of Claim 6 wherein the nucleotide sequence of the fragment comprises all or a portion of the sequence set forth in a member selected from the group consisting of SEQ ID NO:17, 19, 21, 23, 25 and 27.
 - 8. A chimeric gene comprising the nucleic acid fragment of Claim 6 operably linked to suitable regulatory sequences.
 - 9. A transformed host cell comprising the chimeric gene of Claim 8.

- 10. An *Beta vulgaris* sugar transport protein polypeptide comprising all or a substantial portion of the amino acid sequence set forth in a member selected from the group consisting of SEQ ID NO:18, 20, 22, 24, 26 and 28.
- 11. A method of altering the level of expression of a sugar transport protein in a host cell comprising:
 - (a) transforming a host cell with the chimeric gene of any of Claims 3 and 8; and
 - (b) growing the transformed host cell produced in step (a) under conditions that are suitable for expression of the chimeric gene
- wherein expression of the chimeric gene results in production of altered levels of a sugar transport protein in the transformed host cell.
 - 12. A method of obtaining a nucleic acid fragment encoding all or a substantial portion of the amino acid sequence encoding a sugar transport protein comprising:
 - (a) probing a cDNA or genomic library with the nucleic acid fragment of any of Claims 1 and 6;
 - (b) identifying a DNA clone that hybridizes with the nucleic acid fragment of any of Claims 1 and 6;
 - (c) isolating the DNA clone identified in step (b); and
 - (d) sequencing the cDNA or genomic fragment that comprises the clone isolated in step (c)

wherein the sequenced nucleic acid fragment encodes all or a substantial portion of the amino acid sequence encoding a sugar transport protein.

- 13. A method of obtaining a nucleic acid fragment encoding a substantial portion of an amino acid sequence encoding a sugar transport protein comprising:
 - (a) synthesizing an oligonucleotide primer corresponding to a portion of the sequence set forth in any of SEQ ID NOs:1, 3, 5, 7, 9, 11, 13, 15, 17, 19, 21, 23, 25 and 27; and
 - (b) amplifying a cDNA insert present in a cloning vector using the oligonucleotide primer of step (a) and a primer representing sequences of the cloning vector

wherein the amplified nucleic acid fragment encodes a substantial portion of an amino acid sequence encoding a sugar transport protein.

- 14. The product of the method of Claim 12.
- 15. The product of the method of Claim 13.

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